

2/32 Figure 2

	Co	ombination	on Index	
	CD4-I	gG2:T-2	0 Mass	Ratio
Percent Inhibition	25:1 (low)	25:1 (high)	5:1	1:1
95	0.32	0.20	0.22	0.50
90	0.38	0.25	0.27	0.55
85	0.43	0.29	0.30	0.59
80	0.47	0.33	0.34	0.62
75	0.51	0.36	0.37	0.65
79 70	0.54	0.39	0.40	0.67
65	0.58	0.42	0.43	0.70
60	0.61	0.45	0.45	0.73
55	0.65	0.48	0.49	0.75
50	0.69	0.51	0.52	0.78

		T-20			CD4-IgG2	
Percent		Concentration, µg/ml	Dose	Concent	Concentration, µg/ml	Dose
Inhibition	Alone	Inhibition Alone Combination	Reduction	Alone	Alone Combination Reduction	Reduction
66	1.1	0.17	9.9	130	4.3	29
56	0.21	0.044	4.9	19	1.10	17
06	0.10	0.024	4.2	7.8	0.59	13
70	0.025	0.0076	3.3	1.6	0.19	8.4
90	0.011	0.0039	2.8	09.0	0.095	6.3

						4 4 6			T-20	
			PRO 542	42		FA:12				
	•	Concen	oncentration		Concentration	ration,		Concentration,	tration,	
	•	,	M	Dogo		Mu	Dose	u	nM	Dose
Percent	Combination		Mic	Doduction	Alone	Mix	Reduction	Alone	Mix	Mix Reduction Alone Mix Reduction Alone Mix Reduction
Inhibition	Index	Alone	MILA	Negarcion	STORES					
95	0.41	10	10 2.1	4.8	730 2.8	2.8	260	94	19	4.9
)		(,	•	000	, -	150	63	14	4.5
90	0.45	7.0	7.0 1.6	7 .4	370	7.7	27	3	-	<u>.</u>
70	0.47	4.1	0.92	4.5	72	1.2	09	30	8.1	3.7
2 1	. 040	7	990	4.7	28	28 0.87	32	19	19 5.8	3.3
20	0.40	7.1	2							

PRO 542, PA12 and T-20 were used in an approximate 1:1:10 molar concentration ratio.

			2 Odd	(4)		PRO 140	9		T-20	
			FKO 247	74						
		Concen	Concentration,		Concentration,	tration,		Concentration,	ration,	
	•		Mu	Dose	Tu	nM	Dose	nM	4	Dose
Percent	Combination				Alene	Mis	Deduction	Alone	Mix	Reduction
Inhibition	Index	Alone	Mix	Keduction	Alone	MIN	Tron merion			Alone Mix Reduction Alone min accuration and
95	0.40	8.5	8.5 1.9	4.5	19 1.0	1.0	19	140	17	8.2
?		ť	1 2	L V	13	0 77	17	100	13	7.7
90	0.39	7.1	1.3	÷	Ç	;			1	,
7	0.37	5.3	0.87	6.1	7.2	0.46	16	27	7.7	7.4
2 1	7.0	9 7	246 063	7.3	4.9	4.9 0.34	14	40	5.6	7.1
20	0.35	7.0	3							

PRC 542, PRO 140 and T-20 were used in an approximate 2:1:20 molar concentration ratio.

			PRO 542	42		PRO 140	40		T-20	0
		Concentration	tration,		Concentration,	tration,		Concentration,	tration,	
Percent	Percent Combination			Dose	Mu	Z	Dose	\u	nM	Dose
Inhibition	Index	1	Mix	Reduction	Alone	Mix	Alone Mix Reduction Alone Mix Reduction Alone Mix Reduction	Alone	Mix	Reduction
95		61	2.5	24	11.9 0.72	0.72	17	156	22	7.1
06	0.22	32	1.4	23	8.4	0.40	21	96	13	7.4
70	0.19	8.6	0.50	20	4.5	0.14	. 32	40	4.5	8.9
50	0.18	4.7	0.26	18	3.0	3.0 0.074	41	23	2.3	10

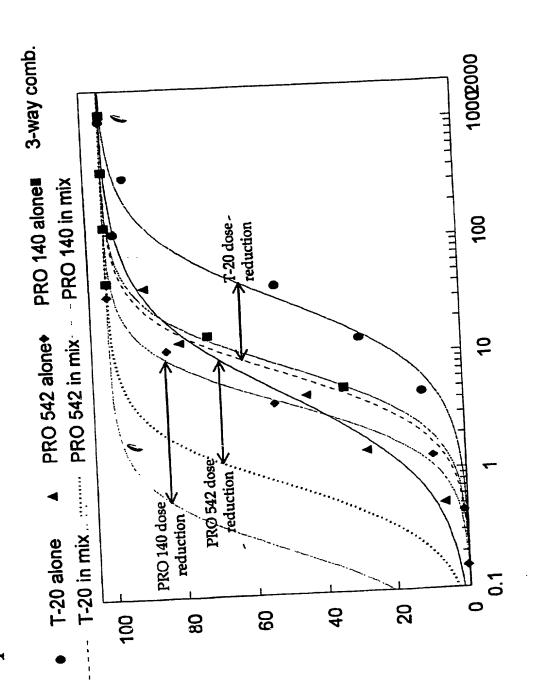
PRO 542, PRO 140 and T-20 were used in an approximate 4:1:30 molar concentration ratio.

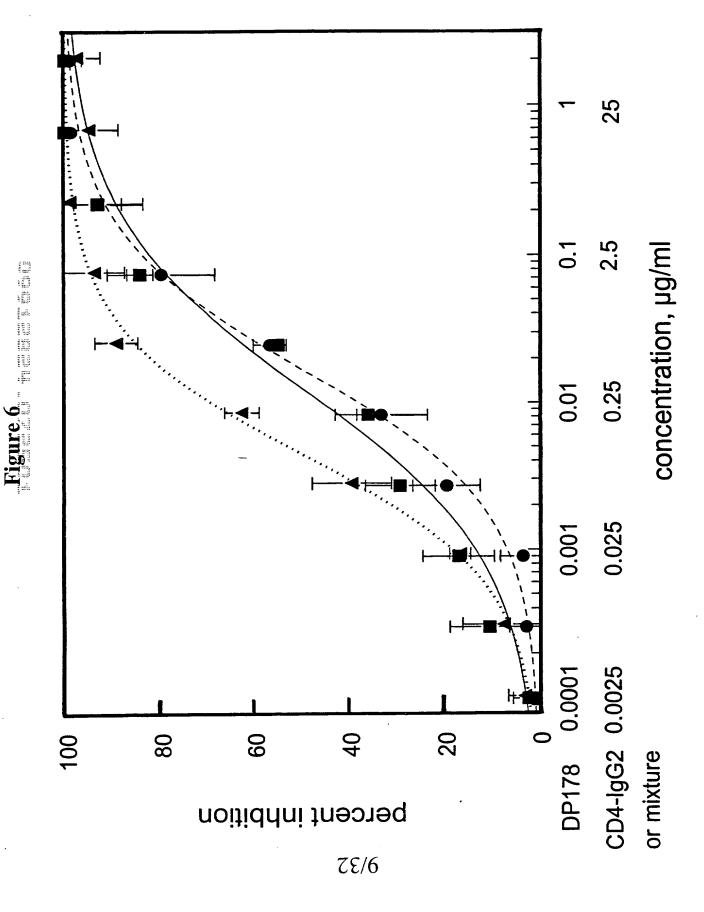
Domont			1	1 TO 140			
Donount Co	•	Concent	Concentration,		Concentration,	tration,	
	Combination	nM	7	Dose	nM	M	Dose
Inhibition	Index	Alone Mix	Mix	Reduction Alone Mix	Alone	Mix	Reduction
96	0.56	12	1.8	6.7	156	55	2.8
06	0.55	8.4	1.1	7.4	96	35	2.7
70	0.55	4.5	0.51	8.8	40	16	2.5
50	0.56	3.0	3.0 0.31	6.6	23	10	2.4

PRO 140 and T-20 were used in an approximate 1:30 molar concentration ratio.

Figure 5 neseron

Triple Combination Synergistically Blocks HIV-1 Entry (I)





	Ö	Combination Index	ion Inde	X
	CD4-1	CD4-IgG2:T-20 Mass Ratio	0 Mass	Ratio
Percent	25:1	25:1		
Inhibition	(low)	(high)	5:1	1:1
95	0.32	0.20	0.22	0.50
06	0.38	0.25	0.27	0.55
85	0.43	0.29	0.30	0.59
80	0.47	0.33	0.34	0.62
75	0.51	0.36	0.37	0.65
70	0.54	0.39	0.40	0.67
65	0.58	0.42	0.43	0.70
. 09	0.61	0.45	0.45	0.73
55	0.65	0.48	0.49	0.75
50	69.0	0.51	0.52	0.78

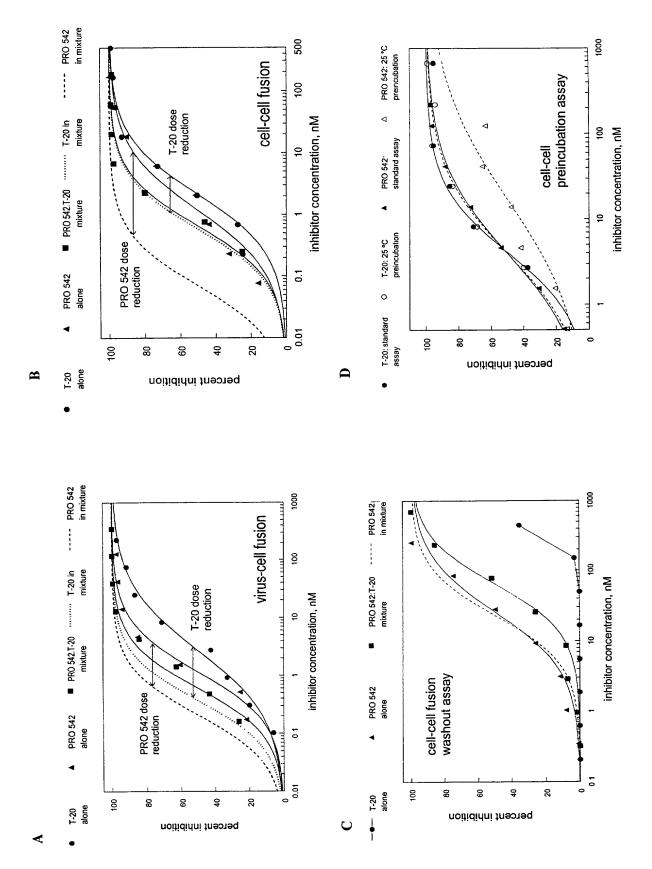
Figure 8

T-20 CD4-IgG2	ration, µg/ml Dose Concentration, µg/ml Dose	Inhibition Alone Combination Reduction Alone Combination Reduction	0.17 6.6 130 4.3 29	0.044 4.9 1.10 1.7	0.024 4.2 7.8 0.59 13	0.0076 3.3 1.6 0.19 8.4	
T-20	itration, µg/ml	Combination R	0.17	0.044	0.024	0.0076	
	Percent Concentral	1 Alone	1.1	0.21	0.10	0.025	
	Percent	Inhibition	66	95	06	70	

Figure 9

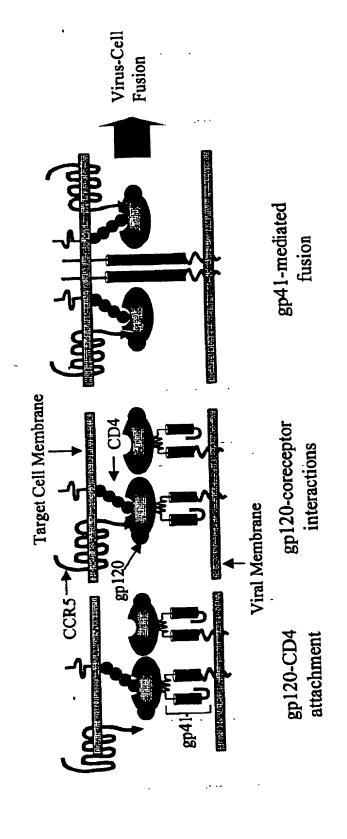
PARTITIONAL OF FORM MACHINE AND PROPERTY IN A STORY AND ADDRESS OF THE PARTITION ADDRESS OF THE PARTITION AND ADDRESS OF THE PARTITION AND ADDRESS OF THE PARTITION ADDRESS OF THE PART	PRO 542:T-20	ANYTHER RESTANDANCES A COMMENT PROGRAMMENT OF THE STANDANCES OF TH	NY IANA TRANSPORTANTANA MARAMPARANTANA TANÀNA TRANSPORTANA	CONTRACTOR OF THE PROPERTY OF	PRO 542	A COMPANY OF THE STATE OF THE S	enterprise and community and the enterprise of t	T-20	engenomen er van damenten. Frankristen en densember van densember
ASSAY (virus)	Molar	rercent Inhihition	Combination	Concentration, nM	ıtion, nM	Dose	Concentration, nM	tion, nM	Dose
(CHIIA)	Ratio	THIRDIAN CHARACTER STREET		Alone	Mix	Reduction	Alone	Mix	Reduction
Virus-cell fusion	1:2	95	0.14	30	2.8	11	120	5.1	24
(JR-FL)		06	0.18	12	1.5	8.0	45	2.6	17
		70	0.29	2.5	0.44	5.7	8.0	0.78	10
		50	0.39	0.92	0.21	4.4	2.7	0.37	7.3
Virus-cell fusion	1:2	95	0.36	65	11	5.9	123	20	6.2
(DH ¹ 23)		06	0.45	20	5.0	4.0	54	8.9	6.1
		70	92.0	2.4	1.2	2.0	12	2.1	5.7
		20	- 1:1	0.64	0.49	1.3	4.8	0.87	5.5
Cell-cell fusion	1:2	95	0.36	35	6.3	5.6	73	11	9.9
(JR-FL)		90	0.43	14	3.2	4.4	34	5.8	5.9
		70	0.61	2.9	0.94	3.1	8.5	1.7	5.0
		20	0.76	1.0	0.43	2.3	3.6	0.78	4.6
Cell-cell fusion	1:10	95	0.27	28	1.4	20	28	12	4.8
(JR-FL)		06	0.28	11	0.55	20	22	4.9	4.5
		70	0.31	2.3	0.11	21	3.8	0.97	3.9
		50	0.34	0.84	0.039	17	1.3	0.35	3.7
Cell-cell fusion	1:50	95	0.33	47	0.84	56	120	37	3.2
(JR-FL)		06	0.34	15	0.30	50	42	13	3.2
		70	0.36	1.8	0.045	40	6.1	2.0	3.0
		50	0.38	0.49	0.014	35	1.8	0.61	3.0
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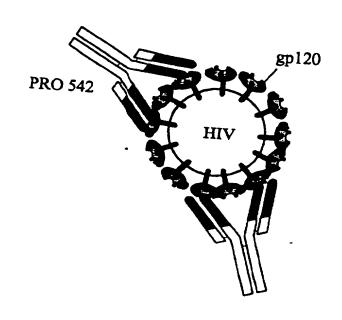


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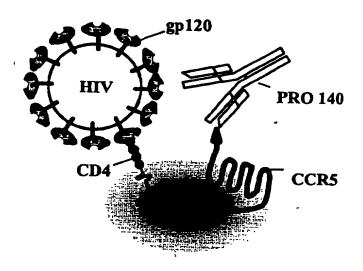
that Provide Promising Targets for Therapy HIV-1 Entry Involves at Least Three Steps



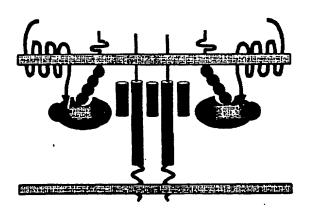
PRO 542 (CD4-IgG2) attachment inhibitor



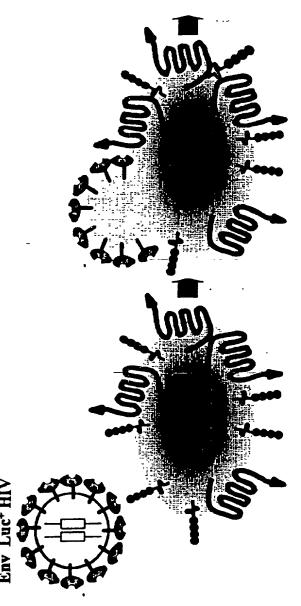
PRO 140 coreceptor inhibitor



T-20 fusion inhibitor



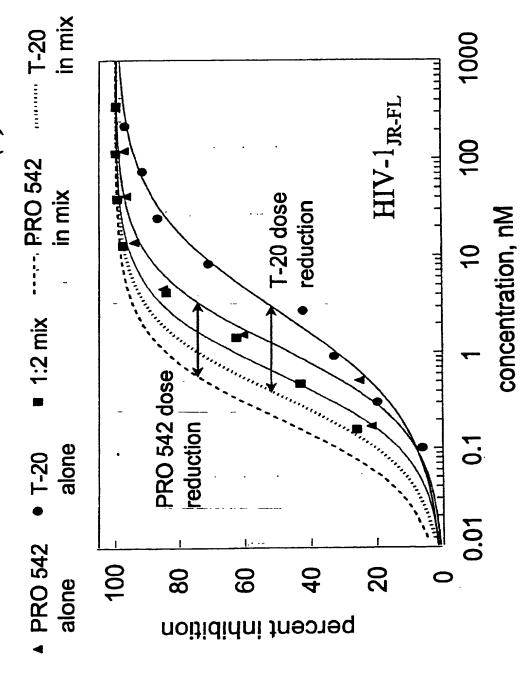
HIV-1 Virus-Cell Fusion Assay



Target Cell

Figure 16

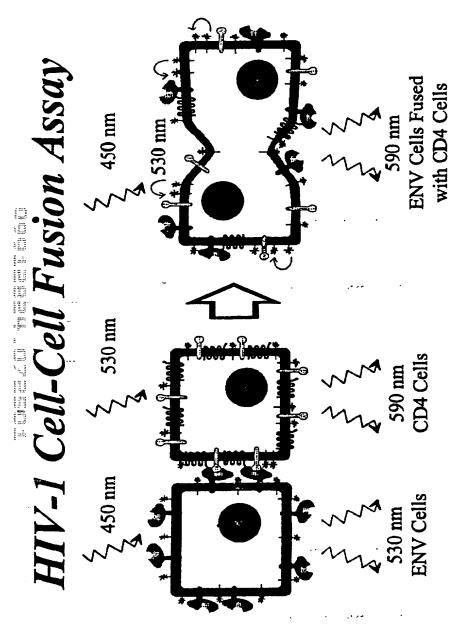
Synergistic Inhibition of Virus-Cell Fusion with PRO 542 and T-20 (I)



Synergistic Inhibition of HIV-1 Virus-Cell-Fusion with PRO 542 and T-20 (II)

Pe	Percent	t Combination Inhibitory Conc., nM	Inhibitory C	onc., nM	Dose Reduction	uction
Inl	Inhibition	Index	PRO 542	T-20	PRO 542 T-20	T-20
JR-FL	95	0.14	30	120	11	24
(R5)	06	0.18	12	45	8.0	17
	20	0.29	2.5	8.0	5.7	10
	20	0.39	0.92	2.7	4.4	7.3
DH123	95	0.36	65	123	5.9	6.2
(R5X4)		0.45	70	54	4.0	6.1
		0.76	2.4	12	2.0	5.7
	20	7:	0.64	4.8	1.3	5.5

PRO 542 and T-20 were used in a 1:2 molar ratio



Synergistic Inhibition of Cell-Cell Fusion with PRO 542 and T-20 (I) Figure 19

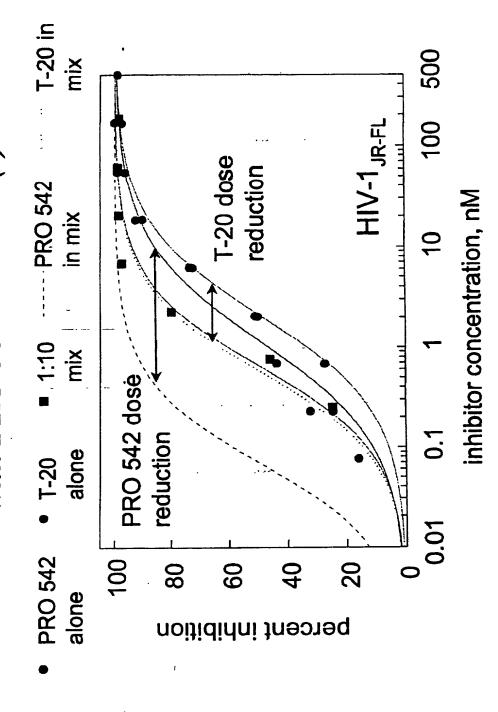


Figure 20

Synergistic Inhibition of HIV-1 Cell-Cell Fusion with PRO 542 and T-20 (II)

Conc.	Conc. Percent (Combination	Combination Inhibitory Conc, nM	onc, nM	Dose Reduction (fold)	tion (fold)
Ratio		Index	PRO 542	T-20	PRO 542	T-20
5	95	0.32	95	47	17	4.9
!	06	0.38	33	22	13	4.2
	20	0.69	3.0	2.5	6.2	2.8
1.10	.•	0.27	28	28	20	8.4
<u>:</u>		0.28	~~	. 22	70	4.5
	20	0.34	0.84	1.3	22	3.7
4.60	95	0.33	47	120	26	3.2
5.		0.34	15	42	20	3.2
	20	0.38	0.49	1.8	35	3.0

Virus: HIV-1_{JR-FL}

PRO 140, PRO 542 and T-20 Triple Combination Figure 21

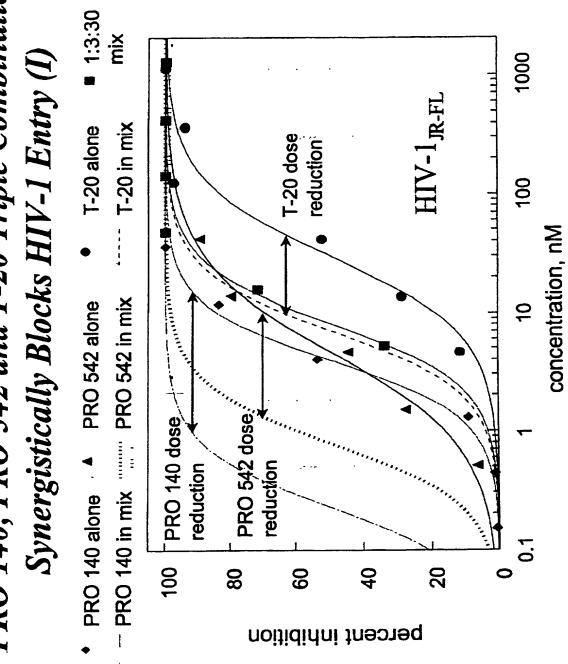


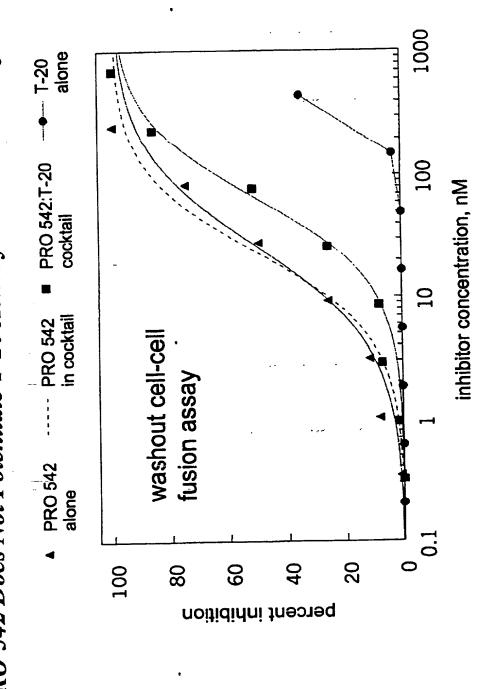
Figure 22 Central of the contract of the contr

PRO 140, PRO 542, T-20 Triple Combination Synergistically Blocks HIV-1 Entry (II)

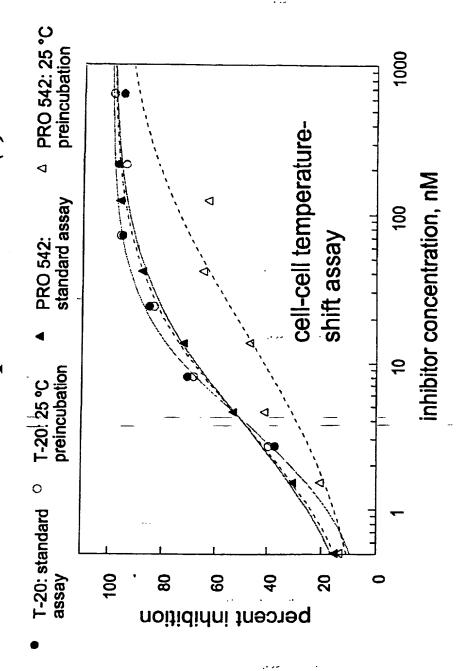
Percent Cor	Combination	Inhibit	Inhibitory Conc, nM	nM	Dose Re	Dose Reduction (fold)	(plo
Inkibition	Index	PRO 140	PRO 542	T-20	PRO 140 PRO 542 T-20 PRO 140 PRO 542 T-20	PRO 542	T-20
95	0.24	24	61	160	17	12	7.1
06	0.22	23	32	96	21	4.8	7.4
20	0.19	70	8.0	40	32	4.5	8.9
20	0.18	18	4.7	23	41	3.0	10

Inhibition of HIV-1_{JR-FL} mediated cell-cell fusion with PRO 140, PRO 542 and T-20 used in a 1:3:30 molar ratio.

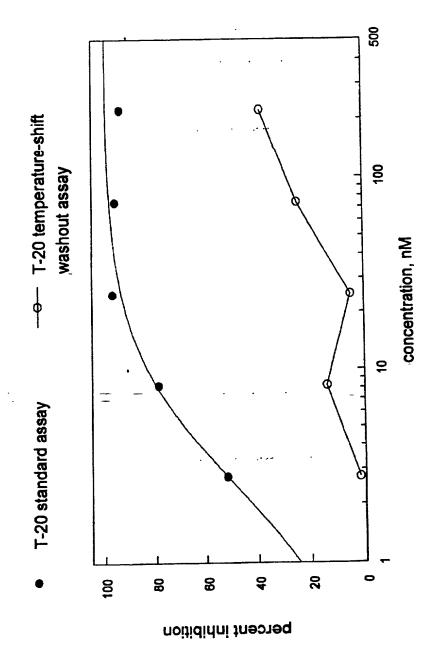
PRO 542 Does Not Potentiate T-20 Activity in the Absence of Coreceptor Figure 23

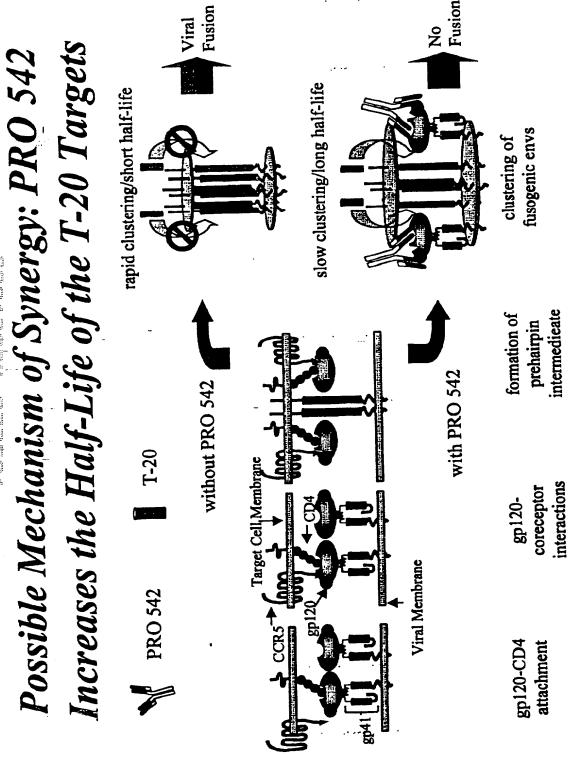


Formation of the Prehairpin Intermediate Requires CD4, Coreceptor and 37 °C(I)



Formation of the Prenairpin Intermediate Requires CD4, Coreceptor and 37 °C (II)





Possible Mechanism of Synergy: PRO 542

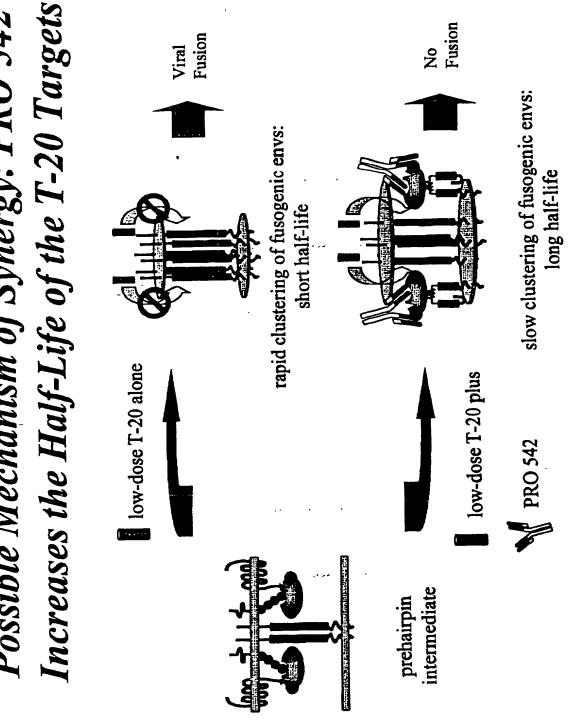
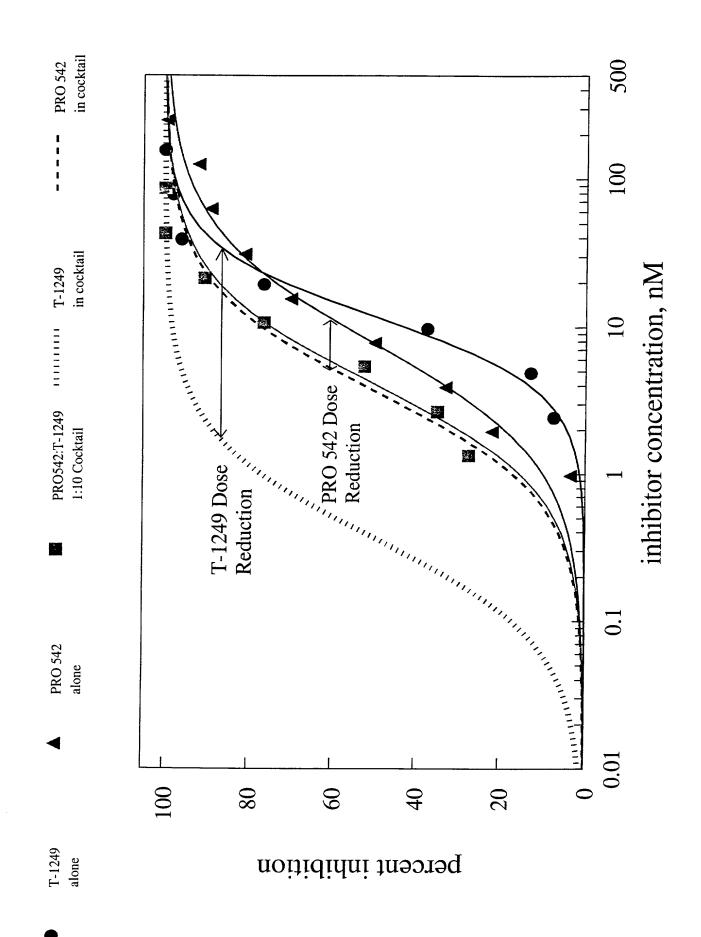


Figure 28 in the second of



						Dose	Dose
Fraction	Dose PRO 542,	Dose PRO 542,	Dose T-1249,	Dose T-1249,	Combination	Reduction	Reduction
Inhibited	nM (alone)	nM (comb)	nM (alone)	nM (comb)	Index	PRO 542	
0.95	87.90	13.58	37.83	1.36	0.20	6.47	27 8G
06.0	48.69	9.52	27.11	0.95	0.24	5 10	28.48
0.85	33.78	7.64	22.06	0.76	0.27	4 42	28.87
0.80	25.65	6.47	18.88	0.65	030	90	20.17
0.75	20.43	5.65	16.61	0.56	0.30	0.00 0.00	20.17
0.70	16.75	5.01	14.85	0000	20.0	3.05	29.42
0.65	12.00	5 5	20:1	00:00	0.34	3.34	29.64
60.0	5.33	4.50	13.41	0.45	0.37	3.11	29.84
0.60	11.81	4.06	12.20	0.41	0.39	2.91	30.03
0.55	10.05	3.68	11.13	0.37	0.41	2.73	30.21
0.50	8.57	3.35	10.18	0.33	0.44	2.56	30.39
) i	